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25 September 1997 (25.09.97)

(54) Title: DETECTION OF NUCLEIC ACID SEQUENCE DIFFERENCES USING THE LIGASE DETECTION REACTION WITH ADDRESSABLE ARRAYS

(57) Abstract

The present invention describes a method for identifying one or more of a plurality of sequences differing by one or more single base changes, insertions, deletions, or translocations in a plurality of target nucleotide sequences. The method includes a ligation phase, a capture phase, and a detection phase. The ligation phase utilizes a ligation detection reaction between one oligonucleotide probe, which has a target sequence-specific portion and an addressable array-specific portion, and a second oligonucleotide probe, having a target sequence-specific portion and a detectable label. After the ligation phase, the capture phase is carried out by hybridizing the ligated oligonucleotide probes to a solid support with an array of immobilized capture oligonucleotides at least some of which are complementary to the addressable array-specific portion. Following completion of the capture phase, a detection phase is carried out to detect the labels of ligated oligonucleotide probes hybridized to the solid support. The ligation phase can be preceded by an amplification process. The present invention also relates to a kit for practicing this method, a method of forming arrays on solid supports, and the supports themselves.

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INTERNATIONAL SEARCH REPORT

International application No. PCT/US97/01535

	ASSIFICATION OF SUBJECT MATTER	,		
IPC(6)	:C07H 21/04, 21/00; C12Q 1/68; C12P 19/34 : 435/6, 91.2; 536/24.3, 25.32	·		
According	to International Patent Classification (IPC) or to be	oth national classification and IPC	•	
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Minimum	documentation searched (classification system follow	wed by classification symbols)		
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0 50				
C. DO	CUMENTS CONSIDERED TO BE RELEVANT			
Category*	Citation of document, with indication, where	appropriate, of the relevant passages	Relevant to claim No.	
Υ	WO 92/10588 A1 (AFFYMAX	TECHNOLOGIES N.V.) 25	1-112 and 120-	
	June 1992, see entire document		147	
Υ	WO 94/11530 A1 (TRUSTEES OF		1-112 and 120-	
	May 1994, see entire document.		147	
Y	NO 00/47400 A4 1747			
7	WO 93/17126 A1 (THE PUB	LIC HEALTH RESEARCH	1-112 and 120-	
	INSTITUTE OF THE CITY OF N	EW YORK) 02 September	147	
	1993, see entire document.	9		
Y,P	US 5,525,464 A(DRMANAC ET	AL \ 11 hans 1000	1 110 1 100	
1	entire document.	AL. IT June 1996, see	1-112 and 120- 147	
ĺ			147	
Y	US 5,412,087/(McGALL ET AL)	02 May 1995, see entire	63, 67-69, 91-	
1	document, especially column 19,	claims 2 and 3.	92, 97, 98,	
l			122, 123, 128	
			and 129	
			2110 120	
Furthe	er documents are listed in the continuation of Box C	See patent family annex.		
	cial categories of cited documents:	"T" Inter document published after the inter	national filing date or priority	
A* door	ament defining the general state of the art which is not considered e of particular relevance	date and not in conflict with the applica principle or theory underlying the inve	tion but cited to understand the ntion	
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INTERNATIONAL SEARCH REPORT

International application No. PCT/US97/01535

Box I Observations where certain claims were found unsearchable (Continuation of item 1 of first sheet)					
This international report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:					
1. Claims Nos.: because they relate to subject matter not required to be searched by this Authority, namely:					
2. Claims Nos.: because they relate to parts of the international application that do not comply with the prescribed requirements to such an extent that no meaningful international search can be carried out, specifically:					
Claims Nos.: because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a). \(\)					
Box II Observations where unity of invention is lacking (Continuation of item 2 of first sheet)					
This International Searching Authority found multiple inventions in this international application, as follows:					
Please See Extra Sheet.					
As all required additional search fees were timely paid by the applicant, this international search report covers all searchable claims.					
2. As all searchable claims could be searched without effort justifying an additional fee, this Authority did not invite payment of any additional fee.					
3. As only some of the required additional search fees were timely paid by the applicant, this international search report covers only those claims for which fees were paid, specifically claims Nos.:					
4. X No required additional search fees were timely paid by the applicant. Consequently, this international search report is restricted to the invention first mentioned in the claims; it is covered by claims Nos.: 1-112 and 120-147					
Remark on Protest The additional search fees were accompanied by the applicant's protest.					
No protest accompanied the payment of additional search fees.					

INTERNATIONAL SEARCH REPORT

International application No. PCT/US97/01535

B. FIELDS SEARCHED

Electronic data bases consulted (Name of data base and where practicable terms used):

USPAT, MEDLINE, BIOSIS, CAPLUS

search terms: array, oligonucleotide probe, labels, ligase, amplification, solid support, capture probe, detect, hybridization

BOX II. OBSERVATIONS WHERE UNITY OF INVENTION WAS LACKING

This ISA found multiple inventions as follows:

This application contains the following inventions or groups of inventions which are not so linked as to form a single inventive concept under PCT Rule 13.1. In order for all inventions to be searched, the appropriate additional search fees must be paid.

Group I, claim(s) 1-112 and 120-147, drawn to a method for identifying one or more sequences differing by one or more single-base.

Group II, claim(s) 113-119, drawn to a device having chambers and valves.

The inventions listed as Groups I do not relate to a single inventive concept under PCT Rule 13.1 because, under PCT Rule 13.2, they lack the same or corresponding special technical features for the following reasons: Group I is drawn to a method for identifying sequences differing by one or more single-base which can be done by hand. Group II is drawn to a device which can be used for protein assay.

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